

=> d his

(FILE 'HOME' ENTERED AT 15:30:42 ON 09 DEC 2008)

FILE 'REGISTRY' ENTERED AT 15:30:58 ON 09 DEC 2008

E TRICARBONYL

L1 72290 S E3

FILE 'CAPLUS' ENTERED AT 15:31:12 ON 09 DEC 2008

L2 30421 S (L1 OR TRICARBONY?)

L3 26829 S L2 AND (PY<=2003)

L4 66 S L3 AND (CANCER? OR TUMOR? OR TUMOUR?)

L5 66 DUP REM L4 (0 DUPLICATES REMOVED)

=> d stat que

L1 72290 SEA FILE=REGISTRY ABB=ON PLU=ON TRICARBONYL/BI

L2 30421 SEA FILE=CAPLUS ABB=ON PLU=ON (L1 OR TRICARBONY?)

L3 26829 SEA FILE=CAPLUS ABB=ON PLU=ON L2 AND (PY<=2003)

L4 66 SEA FILE=CAPLUS ABB=ON PLU=ON L3 AND (CANCER? OR TUMOR? OR  
TUMOUR?)

L5 66 DUP REM L4 (0 DUPLICATES REMOVED)

=> log y

=> d his

(FILE 'HOME' ENTERED AT 22:07:07 ON 09 DEC 2008)

FILE 'REGISTRY' ENTERED AT 22:07:40 ON 09 DEC 2008  
L1           SCREEN 1840 AND 1918 AND 2006 AND 1996  
L2           STRUCTURE uploaded  
L3           QUE L2 AND L1  
L4           0 S L3  
L5           2 S L3 FULL  
L6           ANALYZE L5 1- LC :         4 TERMS

FILE 'CAPLUS' ENTERED AT 22:09:15 ON 09 DEC 2008  
L7           4 S L5  
L8           4 DUP REM L7 (0 DUPLICATES REMOVED)

FILE 'REGISTRY' ENTERED AT 22:09:43 ON 09 DEC 2008  
L9           SCREEN 1838 AND 1918 AND 2008 AND 1993  
L10          STRUCTURE uploaded  
L11          QUE L10 AND L9  
L12          0 S L11  
L13          41 S L11 FULL

FILE 'CAPLUS' ENTERED AT 22:10:29 ON 09 DEC 2008  
L14          13 S L13  
L15          13 DUP REM L14 (0 DUPLICATES REMOVED)

FILE 'REGISTRY' ENTERED AT 22:11:10 ON 09 DEC 2008  
L16          SCREEN 1839 AND 1918 AND 1929 AND 2006 AND 1995  
L17          STRUCTURE uploaded  
L18          QUE L17 AND L16  
L19          0 S L18  
L20          11 S L18 FULL

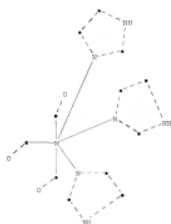
FILE 'CAPLUS' ENTERED AT 22:12:15 ON 09 DEC 2008  
L21          6 S L20  
L22          6 DUP REM L21 (0 DUPLICATES REMOVED)

=> d stat que  
L16          SCR 1839 AND 1918 AND 1929 AND 2006 AND 1995  
L17          STR

\*\*\* STRUCTURE DIAGRAM IS NOT AVAILABLE \*\*\*

Structure attributes must be viewed using STN Express query preparation.  
L20          11 SEA FILE=REGISTRY SSS FUL L17 AND L16  
L21          6 SEA FILE=CAPLUS ABB=ON PLU=ON L20  
L22          6 DUP REM L21 (0 DUPLICATES REMOVED)

=> log y



```

chain nodes :
 1 2 3 4 5 6 7
ring nodes :
 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22
chain bonds :
 1-2 1-3 1-4 1-8 1-9 1-18 2-7 3-6 4-5
ring bonds :
 8-14 8-17 9-10 9-13 10-11 11-12 12-13 14-15 15-16 16-17 18-19 18-22 19-20 20-21
 21-22
exact/norm bonds :
 2-7 3-6 4-5 8-14 8-17 9-10 9-13 10-11 11-12 12-13 14-15 15-16 16-17 18-19 18-22
 19-20 20-21 21-22
exact bonds :
 1-2 1-3 1-4 1-8 1-9 1-18

Match level :
 1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom
 12:Atom 13:Atom 14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom
 22:Atom

```



chain nodes :  
2 3 4 5 6 7 12 13

ring nodes :  
1 8 9 10 11

chain bonds :  
1-2 1-3 1-4 2-7 3-6 4-5 10-13 11-12

ring bonds :  
1-8 1-9 8-11 9-10 10-11

exact/norm bonds :  
1-8 1-9 2-7 3-6 4-5 8-11 9-10 10-11 11-12

exact bonds :  
1-2 1-3 1-4 10-13

Match level :  
1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:Atom 9:Atom 10:Atom 11:Atom  
12:CLASS 13:CLASS



```

chain nodes :
 1  2  3  4  5  6  7  18
ring nodes :
 8  9  10  11  12  13  14  15  16  17
chain bonds :
 1-2  1-3  1-4  1-8  1-9  1-18  2-7  3-6  4-5
ring bonds :
 8-14  8-17  9-10  9-13  10-11  11-12  12-13  14-15  15-16  16-17
exact/norm bonds :
 2-7  3-6  4-5  8-14  8-17  9-10  9-13  10-11  11-12  12-13  14-15  15-16  16-17
exact bonds :
 1-2  1-3  1-4  1-8  1-9  1-18

Match level :
 1:CLASS  2:CLASS  3:CLASS  4:CLASS  5:CLASS  6:CLASS  7:CLASS  8:Atom  9:Atom  10:Atom  11:Atom
 12:Atom  13:Atom  14:Atom  15:Atom  16:Atom  17:Atom  18:CLASS

```